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09/890,811

BB1436

PRELIMINARY AMENDMENT AND RESPONSE TO NOTIFICATION OF MISSING
REQUIREMENT UNDER 35 U.S.C. 371

STATEMENT UNDER 37 CFR 1.821(g) AND 1.825(b)

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Seq List + CRF + 17 Pages Mmc 11-8-01

SEQUENCE LISTING

<110> E.I. du Pont de Nemours and Company

<120> SPFl-Related Transcription Factors

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<150> 60/174325

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 35 40 45
 Leu Phe His Arg Gly Ala Arg Gly Val Pro Lys Phe Lys Ser Ala Gln
 50 55 60
 Pro Pro Ser Leu Pro Ile Ser Pro Pro Pro Met Ser Pro Ser Ser Tyr
 65 70 75 80
 Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser Pro
 85 90 95
 Val Leu Leu His Ser Ser Ser Asn Ile Leu Ala Ser Pro Thr Thr Gly
 100 105 110
 Ala Ile Pro Ala Gln Arg Phe Asp Trp Lys Lys Ala Ala Asp Leu Ile
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 Phe Asp Asp Phe Ser Phe
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 <213> *Oryza sativa*

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 35 40 45
 Arg Gly Gly Ala Arg Val Gly Gly Gly Val Pro Lys Phe Lys Ser Ala
 50 55 60
 Gln Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Val Ser Pro Ser Ser
 65 70 75 80

Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Thr Glu Leu Leu Asp Ser
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 Pro Val Leu Leu Ser Ser Ser His Ile Leu Ala Phe Pro Thr Thr Gly
 100 105 110
 Ala Ile Pro Ala Gln Arg Tyr Asp Trp Lys Ala Ser Ala Asp Leu Ile
 115 120 125
 Ala Ser Gln Gln Asp Asp Ser Arg Gly Asp Phe Ser Phe His Thr Asn
 130 135 140
 Ser Asp Ala Met Ala Ala Gln Pro Ala Ser Phe Pro Ser Phe Lys Glu
 145 150 155 160
 Gln Glu Gln Gln Val Val Glu Ser Ser Lys Asn Gly Ala Ala Ala Ala
 165 170 175
 Ser Ser Asn Lys Ser Gly Gly Gly Gly Asn Asn Lys Leu Glu Asp Gly
 180 185 190
 Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn
 195 200 205
 Pro Arg Ser Tyr Tyr Lys Cys Thr Tyr Asn Gly Cys Ser Met Lys Lys
 210 215 220
 Lys Val Glu Arg Ser Leu Ala Asp Gly Arg Ile Thr Gln Ile Val Tyr
 225 230 235 240
 Lys Gly Ala His Asn His Pro Lys Pro Leu Ser Thr Ala Ala Thr Pro
 245 250 255
 Leu Pro Ala Pro Pro Pro Pro Pro Ala Pro Thr Thr Ser Arg Arg Pro
 260 265 270
 Ala Arg Ala Arg Thr Ser Thr Pro Pro Arg Arg Pro Arg Thr Pro Pro
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 325 330 335
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 340 345 350
 Asn Gly Phe Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys Gly Asn
 355 360 365
 Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Thr Val Gly Cys Pro Val
 370 375 380
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 385 390 395 400

Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Val Arg Pro Arg Arg
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 420 425 430
 Ser Gly Pro Thr Asp Val Ala Ala Ala Gln Gln Gly Pro Tyr Thr Leu
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 Glu Met Leu Pro Asn Pro Ala Gly Leu Tyr Gly Gly Tyr Gly Ala Gly
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 <213> Oryza sativa

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 <213> Oryza sativa

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 Arg Gly Gly Ala Arg Val Gly Gly Gly Val Pro Lys Phe Lys Ser Ala
 50 55 60
 Gln Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Val Ser Pro Ser Ser
 65 70 75 80
 Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Thr Glu Leu Leu Asp Ser
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 Pro Val Leu Leu Ser Ser Ser His Ile Leu Ala Ser Pro Thr Thr Gly
 100 105 110
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 Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn
 195 200 205
 Pro Arg Ser Tyr Tyr Lys Cys Thr Tyr Asn Gly Cys Ser Met Lys Lys
 210 215 220
 Lys Val Glu Arg Ser Leu Ala Asp Gly Arg Ile Thr Gln Ile Val Tyr
 225 230 235 240
 Lys Gly Ala His Asn His Pro Lys Pro Leu Ser Thr Arg Arg Asn Ala
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 Ser Ser Cys Ala Thr Ala Ala Ala Cys Ala Asp Asp Leu Ala Ala Pro
 260 265 270

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 305 310 315 320
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 325 330 335
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 340 345 350
 Asp Gly Phe Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys Gly Asn
 355 360 365
 Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Thr Val Gly Cys Pro Val
 370 375 380
 Arg Lys His Val Glu Arg Ala Ser His Asp Thr Arg Ala Val Ile Thr
 385 390 395 400
 Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Val Gly Arg Gly Gly
 405 410 415
 Gly Gly Gly Arg Ala Pro Ala Pro Ala Pro Pro Thr Ser Gly Ala Ile
 420 425 430
 Arg Pro Ser Ala Val Ala Ala Ala Gln Gln Gly Pro Tyr Thr Leu Glu
 435 440 445
 Met Leu Pro Asn Pro Ala Gly Leu Tyr Gly Gly Tyr Gly Ala Gly Ala
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 <213> Glycine max

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      35                      40                      45

Gly Leu Ser Glu Arg Thr Gly Ser Gly Val Pro Lys Phe Lys Ser Thr
      50                      55                      60

Pro Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Ile Ser Pro Ser Ser
      65                      70                      75                      80

Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser
      85                      90                      95

Pro Val Leu Leu Asn Ser Ser Asn Ile Leu Pro Ser Pro Thr Thr Gly
      100                     105                     110

Ala Phe Val Ala Gln Ser Phe Asn Trp Lys Ser Ser Ser Gly Gly Asn
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Gln Thr Arg Ser Gly Pro Pro Ala Ser Ser Thr Ala Thr Tyr Gln Ser
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Ser Asn Val Thr Val Gln Thr Gln Gln Pro Trp Ser Phe Gln Glu Ala
 165 170 175
 Thr Lys Gln Asp Asn Phe Ser Ser Gly Lys Gly Met Met Lys Thr Glu
 180 185 190
 Asn Ser Ser Ser Met Gln Ser Phe Ser Pro Glu Ile Ala Ser Val Gln
 195 200 205
 Thr Asn His Ser Asn Gly Phe Gln Ser Asp Tyr Gly Asn Tyr Pro Pro
 210 215 220
 Gln Ser Gln Thr Leu Ser Arg Arg Ser Asp Asp Gly Tyr Asn Trp Arg
 225 230 235 240
 Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn Pro Arg Ser Tyr
 245 250 255
 Tyr Lys Cys Thr Tyr Pro Asn Cys Pro Thr Lys Lys Lys Val Glu Arg
 260 265 270
 Ser Leu Asp Gly Gln Ile Thr Glu Ile Val Tyr Lys Gly Thr His Asn
 275 280 285
 His Pro Lys Pro Gln Asn Thr Arg Arg Asn Ser Ser Asn Ser Ser Ser
 290 295 300
 Leu Ala Ile Pro His Ser Asn Ser Ile Arg Thr Glu Ile Pro Asp Gln
 305 310 315 320
 Ser Tyr Ala Thr His Gly Ser Gly Gln Met Asp Ser Ala Ala Thr Pro
 325 330 335
 Glu Asn Ser Ser Ile Ser Ile Gly Asp Asp Asp Phe Glu Gln Ser Ser
 340 345 350
 Gln Lys Cys Lys Ser Gly Gly Asp Glu Tyr Asp Glu Asp Glu Pro Asp
 355 360 365
 Ala Lys Arg Trp Lys Ile Glu Gly Glu Asn Glu Gly Met Ser Ala Pro
 370 375 380
 Gly Ser Arg Thr Val Arg Glu Pro Arg Val Val Val Gln Thr Thr Ser
 385 390 395 400
 Asp Ile Asp Ile Leu Asp Asp Gly Tyr Arg Trp Arg Lys Tyr Gly Gln
 405 410 415
 Lys Val Val Lys Gly Asn Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr
 420 425 430
 His Pro Gly Cys Pro Val Arg Lys His Val Glu Arg Ala Ser His Asp
 435 440 445
 Leu Arg Ala Val Ile Thr Thr Tyr Glu Gly Lys His Asn His Asp Val
 450 455 460
 Pro Ala Ala Arg Gly Ser Gly Ser His Ser Val Asn Arg Pro Met Pro
 465 470 475 480

Asn Asn Ala Ser Asn His Thr Asn Thr Ala Ala Thr Ser Val Arg Leu
 485 490 495
 Leu Pro Val Ile His Gln Ser Asp Asn Ser Leu Gln Asn Gln Arg Ser
 500 505 510
 Gln Ala Pro Pro Glu Gly Gln Ser Pro Phe Thr Leu Glu Met Leu Gln
 515 520 525
 Ser Pro Gly Ser Phe Gly Phe Ser Gly Phe Gly Asn Pro Met Gln Ser
 530 535 540
 Tyr Val Asn Gln Gln Gln Leu Ser Asp Asn Val Phe Ser Ser Arg Thr
 545 550 555 560
 Lys Glu Glu Pro Arg Asp Asp Met Phe Leu Glu Ser Leu Leu Cys
 565 570 575

<210> 11
 <211> 2158
 <212> DNA
 <213> Triticum aestivum

<400> 11
 gcacgagccg caccgcgcgc atggccgatt cgccaaaccc tagctccggg gacctcccct 60
 cagccgcgcg gagctcgccc gagaagccgt accccgcgga tcgacgcgtc ggggcgctcg 120
 ccggcgcggg cgcgaggtac aaggccatgt ccccggcgcg gctgccgata tcgcgcgagc 180
 cctgcctcac catccccgcc ggcttcagcc cctccgccct cctcgactcc cccgtgctcc 240
 tcaccaactt caaggttgaa ccttcaccaa caactggtag tctgagcatg gctgcaatta 300
 tgcaacaagag tgctcatcca gacatactgc cttcgccacg ggataagtct attcgagccc 360
 atgaagatgg gggttctagg gattttgaat tcaagcctca tctgaattcg tcttctcaat 420
 cactggctcc tgctatgagt gatctaaaaa aacacgagca ttctatgcaa aatcagagta 480
 tgaatcccag ctcatcatct agcaatatgg tgaatgaaaa cagacctccc tgttcacgcg 540
 agtcaagtct tacagtgaat gtaagtgtct cgaaccaacc tgttggaatg gttggtttga 600
 ctgacaacat gcctgctgaa gttggtacat ctgagccgca gcagatgaat agttctgaca 660
 atgccatgca agagccgcag tctgaaaatg ttgctgacaa gtcagcagat gatggctaca 720
 actggcgcaa atatgggcag aagcatgtca agggaagtga aaaccctaga agttattaca 780
 agtgcacaca tcctaattgt gaagtaaaaa agctattgga gcgtgcgggt gatggtctga 840
 tcacggaagt tgtctataag gggcgccata atcatcctaa gccccagcct aataggaggt 900
 tagctgggtg tgcaagttct tcgaaccagg gtgaagaacg atatgatggg gcggcagctg 960
 ctgatgataa atcttccaat gctcttagca accttgctaa tccggtaaat tcgcctggca 1020
 tggttgagcc tgttccagtt tcagttagtg atgatgacat agatgctgga ggtggaagac 1080
 cctaccctgg ggatgatgct acagaggagg atttagagtc gaaacgcagg aaaatggagt 1140
 ctgcaggatg tgatgctgct ctgatgggta aacctaaccg tgagccccgt gttgtcggtc 1200
 agactgtaag tgaggttgac atcttggatg atgggtatcg ttggcggaaa tatggacaga 1260
 aagttgtcaa aggaaacccc aatccacgga gttactacaa atgcacaagc acaggatgcc 1320
 ctgtgaggaa gcatgttgag agagcatcgc acgaccta atcagtgata acaacgtatg 1380
 aaggaaaaca taacctgaa gtccctgctg cgaggaatgc aacctatgag atgtccgcgc 1440
 ctcccatgaa gaatgtcgtg catcagatta acagcagtat gcccagcagc attggcggca 1500
 tgatgagagc atgtgaagcc aggaacttca gcaaccaata ttctcaagcc gctgaaaccg 1560
 acaatgtcag tcttgacctt ggtgttgagg tcagcccga ccacagcgat gccacaaacc 1620
 aaatgcagtc ttcaggctct gatcagatgc agtaccagat gcaatccatg gcttcgatgt 1680
 acggcaacat gagacatcca tcatcaatgg cagtgccaac ggtacaagga aactctgctg 1740
 gccgcatgta tggttccaga gaagagaaaag gtaacgaagg gtttactttc agagccacac 1800
 cgatggacca ttcagctaac ctatgctata gcggtgctgg gaacttggtc atgggtccat 1860
 gagaggaatg atgagagtgt cagcaaatgc ttatagctcc atgaatcata tattacaac 1920
 aatgcttttg taacgacaat ctcttcagca agattcttaa ttgtgtatcg gttacaagtc 1980
 agttcagcca gaggcaagta agctataagc tatacctgga ggactgcagc aaatgcgcat 2040

gtgtcttttt aggcgcggaa aaggcccctg ctgtatgtag cgctgcagac ctacattcgt 2100
 tgtacagcga acctaatatg attaattaat tagattatga gaatttggtt taaaaaaaa 2158

<210> 12
 <211> 619
 <212> PRT
 <213> Triticum aestivum

<400> 12
 Thr Ser Arg Thr Ala Pro Met Ala Asp Ser Pro Asn Pro Ser Ser Gly
 1 5 10 15
 Asp Leu Pro Ser Ala Ala Gly Ser Ser Pro Glu Lys Pro Tyr Pro Ala
 20 25 30
 Asp Arg Arg Val Ala Ala Leu Ala Gly Ala Gly Ala Arg Tyr Lys Ala
 35 40 45
 Met Ser Pro Ala Arg Leu Pro Ile Ser Arg Glu Pro Cys Leu Thr Ile
 50 55 60
 Pro Ala Gly Phe Ser Pro Ser Ala Leu Leu Asp Ser Pro Val Leu Leu
 65 70 75 80
 Thr Asn Phe Lys Val Glu Pro Ser Pro Thr Thr Gly Ser Leu Ser Met
 85 90 95
 Ala Ala Ile Met His Lys Ser Ala His Pro Asp Ile Leu Pro Ser Pro
 100 105 110
 Arg Asp Lys Ser Ile Arg Ala His Glu Asp Gly Gly Ser Arg Asp Phe
 115 120 125
 Glu Phe Lys Pro His Leu Asn Ser Ser Ser Gln Ser Leu Ala Pro Ala
 130 135 140
 Met Ser Asp Leu Lys Lys His Glu His Ser Met Gln Asn Gln Ser Met
 145 150 155 160
 Asn Pro Ser Ser Ser Ser Ser Asn Met Val Asn Glu Asn Arg Pro Pro
 165 170 175
 Cys Ser Arg Glu Ser Ser Leu Thr Val Asn Val Ser Ala Pro Asn Gln
 180 185 190
 Pro Val Gly Met Val Gly Leu Thr Asp Asn Met Pro Ala Glu Val Gly
 195 200 205
 Thr Ser Glu Pro Gln Gln Met Asn Ser Ser Asp Asn Ala Met Gln Glu
 210 215 220
 Pro Gln Ser Glu Asn Val Ala Asp Lys Ser Ala Asp Asp Gly Tyr Asn
 225 230 235 240
 Trp Arg Lys Tyr Gly Gln Lys His Val Lys Gly Ser Glu Asn Pro Arg
 245 250 255
 Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys Glu Val Lys Lys Leu Leu
 260 265 270

Glu Arg Ala Val Asp Gly Leu Ile Thr Glu Val Val Tyr Lys Gly Arg
 275 280 285
 His Asn His Pro Lys Pro Gln Pro Asn Arg Arg Leu Ala Gly Gly Ala
 290 295 300
 Val Pro Ser Asn Gln Gly Glu Glu Arg Tyr Asp Gly Ala Ala Ala Ala
 305 310 315 320
 Asp Asp Lys Ser Ser Asn Ala Leu Ser Asn Leu Ala Asn Pro Val Asn
 325 330 335
 Ser Pro Gly Met Val Glu Pro Val Pro Val Ser Val Ser Asp Asp Asp
 340 345 350
 Ile Asp Ala Gly Gly Gly Arg Pro Tyr Pro Gly Asp Asp Ala Thr Glu
 355 360 365
 Glu Asp Leu Glu Ser Lys Arg Arg Lys Met Glu Ser Ala Gly Ile Asp
 370 375 380
 Ala Ala Leu Met Gly Lys Pro Asn Arg Glu Pro Arg Val Val Val Gln
 385 390 395 400
 Thr Val Ser Glu Val Asp Ile Leu Asp Asp Gly Tyr Arg Trp Arg Lys
 405 410 415
 Tyr Gly Gln Lys Val Val Lys Gly Asn Pro Asn Pro Arg Ser Tyr Tyr
 420 425 430
 Lys Cys Thr Ser Thr Gly Cys Pro Val Arg Lys His Val Glu Arg Ala
 435 440 445
 Ser His Asp Pro Lys Ser Val Ile Thr Thr Tyr Glu Gly Lys His Asn
 450 455 460
 His Glu Val Pro Ala Ala Arg Asn Ala Thr His Glu Met Ser Ala Pro
 465 470 475 480
 Pro Met Lys Asn Val Val His Gln Ile Asn Ser Ser Met Pro Ser Ser
 485 490 495
 Ile Gly Gly Met Met Arg Ala Cys Glu Ala Arg Asn Phe Ser Asn Gln
 500 505 510
 Tyr Ser Gln Ala Ala Glu Thr Asp Asn Val Ser Leu Asp Leu Gly Val
 515 520 525
 Gly Ile Ser Pro Asn His Ser Asp Ala Thr Asn Gln Met Gln Ser Ser
 530 535 540
 Gly Pro Asp Gln Met Gln Tyr Gln Met Gln Ser Met Ala Ser Met Tyr
 545 550 555 560
 Gly Asn Met Arg His Pro Ser Ser Met Ala Val Pro Thr Val Gln Gly
 565 570 575
 Asn Ser Ala Gly Arg Met Tyr Gly Ser Arg Glu Glu Lys Gly Asn Glu
 580 585 590

Gly Phe Thr Phe Arg Ala Thr Pro Met Asp His Ser Ala Asn Leu Cys
595 600 605

Tyr Ser Gly Ala Gly Asn Leu Val Met Gly Pro
610 615

<210> 13

<211> 549

<212> PRT

<213> Ipomoea batatas

<400> 13

Met Ala Ala Ser Ser Gly Thr Ile Asp Ala Pro Thr Ala Ser Ser Ser
1 5 10 15

Phe Ser Phe Ser Thr Ala Ser Ser Phe Met Ser Ser Phe Thr Asp Leu
20 25 30

Leu Ala Ser Asp Ala Tyr Ser Gly Gly Ser Val Ser Arg Gly Leu Gly
35 40 45

Asp Arg Ile Ala Glu Arg Thr Gly Ser Gly Val Pro Lys Phe Lys Ser
50 55 60

Leu Pro Pro Pro Ser Leu Pro Leu Ser Ser Pro Ala Val Ser Pro Ser
65 70 75 80

Ser Tyr Phe Ala Phe Pro Pro Gly Leu Ser Pro Ser Glu Leu Leu Asp
85 90 95

Ser Pro Val Leu Leu Ser Ser Ser Asn Ile Leu Pro Ser Pro Thr Thr
100 105 110

Gly Thr Phe Pro Ala Gln Thr Phe Asn Trp Lys Asn Asp Ser Asn Ala
115 120 125

Ser Gln Glu Asp Val Lys Gln Glu Glu Lys Gly Tyr Pro Asp Phe Ser
130 135 140

Phe Gln Thr Asn Ser Ala Ser Met Thr Leu Asn Tyr Glu Asp Ser Lys
145 150 155 160

Arg Lys Asp Glu Leu Asn Ser Leu Gln Ser Leu Pro Pro Val Thr Thr
165 170 175

Ser Thr Gln Met Ser Ser Gln Asn Asn Gly Gly Ser Tyr Ser Glu Tyr
180 185 190

Asn Asn Gln Cys Cys Pro Pro Ser Gln Thr Leu Arg Glu Gln Arg Arg
195 200 205

Ser Asp Asp Gly Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys
210 215 220

Gly Ser Glu Asn Pro Arg Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys
225 230 235 240

Pro Thr Lys Lys Lys Val Glu Arg Ala Leu Asp Gly Gln Ile Thr Glu
 245 250 255
 Ile Val Tyr Lys Gly Ala His Asn His Pro Lys Pro Gln Ser Thr Arg
 260 265 270
 Arg Ser Ser Ser Ser Thr Ala Ser Ser Ala Ser Thr Leu Ala Ala Gln
 275 280 285
 Ser Tyr Asn Ala Pro Ala Ser Asp Val Pro Asp Gln Ser Tyr Trp Ser
 290 295 300
 Asn Gly Asn Gly Gln Met Asp Ser Val Ala Thr Pro Glu Asn Ser Ser
 305 310 315 320
 Ile Ser Val Gly Asp Asp Glu Phe Glu Gln Ser Ser Gln Lys Arg Glu
 325 330 335
 Ser Gly Gly Asp Glu Phe Asp Glu Asp Glu Pro Asp Ala Lys Arg Trp
 340 345 350
 Lys Val Glu Asn Glu Ser Glu Gly Val Ser Ala Gln Gly Ser Arg Thr
 355 360 365
 Val Arg Glu Pro Arg Val Val Val Gln Thr Thr Ser Asp Ile Asp Ile
 370 375 380
 Leu Asp Asp Gly Tyr Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys
 385 390 395 400
 Gly Asn Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Ser Gln Gly Cys
 405 410 415
 Pro Val Arg Lys His Val Glu Arg Ala Ser His Asp Ile Arg Ser Val
 420 425 430
 Ile Thr Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Ala Ala Arg
 435 440 445
 Gly Ser Gly Ser His Gly Leu Asn Arg Gly Ala Asn Pro Asn Asn Asn
 450 455 460
 Ala Ala Met Ala Met Ala Ile Arg Pro Ser Thr Met Ser Leu Gln Ser
 465 470 475 480
 Asn Tyr Pro Ile Pro Ile Pro Ser Thr Arg Pro Met Gln Gln Gly Glu
 485 490 495
 Gly Gln Ala Pro Tyr Glu Met Leu Gln Gly Ser Gly Gly Phe Gly Tyr
 500 505 510
 Ser Gly Phe Gly Asn Pro Met Asn Ala Tyr Ala Asn Gln Ile Gln Asp
 515 520 525
 Asn Ala Phe Ser Arg Ala Lys Glu Glu Pro Arg Asp Asp Leu Phe Leu
 530 535 540
 Asp Thr Leu Leu Ala
 545

<210> 14
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Synthetic oligonucleotide

<400> 14
gaaaaattca tcagtggatc cttatttgac ctgtct 36